SEQUENCE LISTING

<110> Leukotech A/S

<120> Pro-inflammatory and anti-inflammatory antibodies against the heparin
binding protein (HBP)

<130> P818 PC00

<160> 589

<170> PatentIn version 3.1

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<213> Homo sapiens

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Gly Val Ser Thr Val Val Leu Gly Ala Tyr Asp Leu Arg Arg Glu 50 55 60

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Thr Gly Val Leu Thr Arg Arg Gly Gly Ile Cys Asn Gly Asp Gly Gly 165 170 175

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Leu Pro Leu Pro

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Gln Glu Phe Pro
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Pro Phe Leu Ala
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<223> Peptide fragment: amino acid residues 26-29 of pHBP
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Cys Ala Gly Ala
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<210> 241
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His Pro Arg Phe
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Asp Pro Arg Gln
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Pro Arg Gln Asn
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Val Leu Leu Leu
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Glu Ala Arg Leu
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Ala Arg Leu Thr
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Ser Val Ala Leu
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Leu Val Pro Leu
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Val Pro Leu Pro
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Gly Trp Gly Thr
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Trp Gly Thr Gln
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Gly Thr Gln Arg
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Gln Arg Leu Arg
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Arg Arg Leu Phe
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Arg Leu Phe Ser
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<210> 316
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Phe Ser Arg Phe
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Phe Pro Arg Val
<210> 318
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Arg Val Leu Asn
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Cys Leu Pro Arg
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Leu Pro Arg Asp
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Arg Asp Met Cys
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Cys Ile Gly Val
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Gly Arg Ile Ser
<210> 343
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Arg Ile Ser Gln
<210> 344
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Ile Ser Gln Gly
<210> 345
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Ser Gln Gly Asp
<210> 346 ·
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Gln Gly Asp Arg
<210> 347
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Asp Arg Gly Thr
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Val Cys Asn Gly
<210> 352
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<223> Peptide fragment: amino acid residues 179-182 of pHBP
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Cys Asn Gly Leu
<210> 353
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Asn Gly Leu Ala
<210> 354
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<223> Peptide fragment: amino acid residues181-184 of pHBP
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Gly Leu Ala Gln
<210> 355
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<223> Peptide fragment: amino acid residues 182-185 of pHBP
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Leu Ala Gln Gly
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<210> 356
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<400> 356
Ala Gln Gly Val
<210> 357
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<223> Peptide fragment: amino acid residues 184-187 of pHBP
<400> 357 `
Gln Gly Val Ala
<210> 358
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<212> PRT
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<223> Peptide fragment: amino acid residues 187-190 of pHBP
<400> 358
Ala Ser Phe Leu
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Ser Phe Leu Arg
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Phe Leu Arg Arg
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Gln Lys Gln Gly
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<400> 362
Arg Arg Arg Phe
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<223> Peptide fragment: amino acid residues 192-195 of pHBP
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Arg Arg Phe Arg
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Phe Arg Arg Ser
<210> 366
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Ser Ser Gly Phe
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<210> 370
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Lys Gln Gly Arg
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Leu Phe Arg Asn
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Ile Asp Ser Val
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Asp Ser Val Leu
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Arg Arg Ala Arg
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Arg Ala Arg Pro
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Arg Pro His Ala
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Gly His Phe Cys
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Cys Gly Ala Thr
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Gly Ala Thr Leu
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<210> 407
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Ala His Cys Val
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<210> 419
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Val Ala Asn Val
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Ala Asn Val Asn
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Val Asn Val Arg
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Val Arg Ala Val
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Gly Ala His Asn
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Glu Pro Thr Arg
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Asp Gly Tyr Asp
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Tyr Asp Pro Val
<210> 454
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Pro Val Asn Leu
<210> 456
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<210> 466
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Asn Gly Ser Ala
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Gly Ser Ala Thr
<210> 470
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Ser Ala Thr Ile
<210> 471
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Ala Thr Ile Asn
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Ser Val Ala Leu
<210> 477
<211> 4
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<223> Peptide fragment: amino acid residues 104-107 of hNLE
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Val Ala Leu Val
<210> 478
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<223> Peptide fragment: amino acid residues 105-108 of hNLE
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Ala Leu Val Pro
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Leu Val Pro Leu
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Pro Leu Pro Ala
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Gln Gly Arg Arg
<210> 486
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Pro Arg Phe Val Leu Thr Ala Ala Ser Cys Phe Arg Gly Lys Asn Ser 35 40 45

Gly Ser Ala Ser Val Val Leu Gly Ala Tyr Asp Leu Arg Gln Glu 50 55 60

Gln Ser Arg Gln Thr Phe Ser Ile Arg Ser Ile Ser Gln Asn Gly Tyr 65 70 75 80

Asp Pro Arg Gln Asn Leu Asn Asp Val Leu Leu Gln Leu Asp Arg 85 90 95

Glu Ala Arg Leu Thr Pro Ser Val Ala Leu Val Pro Leu Pro Pro Gln
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Asn Ala Thr Val Glu Ala Gly Thr Asn Cys Gln Val Glu Ala Gly Trp
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Gly Thr Gln Arg Leu Arg Arg Leu Phe Ser Arg Phe Pro Arg Val Leu 130 135 140

Asn Val Thr Val Thr Ser Asn Pro Cys Leu Pro Arg Asp Met Cys Ile 145 150 155 160 Gly Val Phe Ser Arg Arg Gly Arg Ile Ser Gln Gly Asp Arg Gly Thr
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Pro Leu Val Cys Asn Gly Leu Ala Gln Gly Val Ala Ser Phe Leu Arg 180 185 190

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Leu Arg Gly Gly His Phe Cys Gly Ala Thr Leu Ile Ala Pro Asn Phe 50 55 60

Val Met Ser Ala Ala His Cys Val Ala Asn Val Asn Val Arg Ala Val 65 70 75 80

Arg Val Val Leu Gly Ala His Asn Leu Ser Arg Arg Glu Pro Thr Arg 85 90 95

Gln Val Phe Ala Val Gln Arg Ile Phe Glu Asn Gly Tyr Asp Pro Val

Asn Leu Leu Asn Asp Ile Val Ile Leu Gln Leu Asn Gly Ser Ala Thr Ile Asn Ala Asn Val Gln Val Ala Gln Leu Pro Ala Gln Gly Arg Arg Leu Gly Asn Gly Val Gln Cys Leu Ala Met Gly Trp Gly Leu Leu Gly Arg Asn Arg Gly Ile Ala Ser Val Leu Gln Glu Leu Asn Val Thr Val Val Thr Ser Leu Cys Arg Arg Ser Asn Val Cys Thr Leu Val Arg Gly Arg Gln Ala Gly Val Cys Phe Gly Asp Ser Gly Ser Pro Leu Val Cys Asn Gly Leu Ile His Gly Ile Ala Ser Phe Val Arg Gly Gly Cys Ala Ser Gly Leu Tyr Pro Asp Ala Phe Ala Pro Val Ala Gln Phe Val Asn Trp Ile Asp Ser Ile Ile Gln Arg Ser Glu Asp Asn Pro Cys Pro His Pro Arg Asp Pro Asp Pro Ala Ser Arg Thr His

